Regulatory Mechanisms of Growth Hormone Secretion Are Sexually Dimorphic

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Abstract

Sexually dimorphic growth hormone (GH) secretory pattern is important in the determination of gender-specific patterns of growth and metabolism in rats. Whether GH secretion in humans is also sexually dimorphic and the neuroendocrine mechanisms governing this potential difference are not fully established. We have compared pulsatile GH secretion profiles in young men and women in the baseline state and during a continuous intravenous infusion of recombinant human insulin-like growth factor I (rhIGF-I). During the baseline study, men had large nocturnal GH pulses and relatively small pulses during the rest of the day. In contrast, women had more continuous GH secretion and more frequent GH pulses that were of more uniform size. The infusion of rhIGF-I (10 μg/kg/h) potently suppressed both spontaneous and growth hormone–releasing hormone (GHRH)-induced GH secretion in men. In women, however, rhIGF-I had less effect on pulsatile GH secretion and did not suppress the GH response to GHRH. These data demonstrate the existence of sexual dimorphism in the regulatory mechanisms involved in GH secretion in humans. The persistence of GH responses to GHRH in women suggests that negative feedback by IGF-I might be expressed, in part, through suppression of hypothalamic GHRH. (J. Clin. Invest. 1998. 102:153–164.) Key words: pituitary • hypothalamus • human • insulin-like growth factor I • thyroid-stimulating hormone

Introduction

The sexual dimorphism of growth hormone (GH) secretion in rats has been extensively studied. High amplitude GH pulses and low interpulse GH concentrations characterizes GH profiles in males, whereas female rats have less regular pulses and higher interpulse GH levels (1). These differences have physiological importance and are responsible for the differential expression of sex-specific, hepatic P450 drug-metabolizing enzymes (2) and for male–female differences in growth rate (3). Potential differences in GH secretion in humans have been less well studied.

Women were found to have higher average daily GH concentrations by some (4–6) but not all (7) investigators. A recent study using deconvolution analysis to estimate GH secretion found twofold higher GH secretion rates in middle-aged women compared with men of a similar age (6); in addition, women appear to secrete GH with more “process irregularity” than men (8). Whether there is a gender-based difference in pituitary sensitivity to GH-releasing hormone (GHRH) is unclear; it has been reported that women have greater (9), smaller (10), or equivalent (11, 12) GH responses to GHRH compared with men. It is also uncertain whether there is sexual dimorphism in the GH responses to provocative pharmacologic testing. Gender differences during testing with pyridostigmine or insulin were observed by some (13, 14) but not all investigators (12). In addition, one study convincingly demonstrated that GH suppression after ingestion of oral glucose was less complete in women than in men (15). Potential mechanisms underlying gender differences in GH secretion have been recently reviewed (16) and might include differences in GH clearance, in GHRH or somatostatin (SRIH) secretion, or in negative feedback at the level of the pituitary or hypothalamus by GH or insulin-like growth factor I (IGF-I).

Accurate analysis of potential sex-specific patterns of GH secretion has been limited by study design or by methodology. Some of the daily GH secretion data were obtained using infrequent blood sampling (5) or relatively insensitive GH assays that did not allow accurate resolution of GH pulsatility in the low assay range (4). One study used a continuous blood draw technique, so potential gender-related differences in GH secretion pattern could not be ascertained (7). In addition, heterogeneous subject populations, the interdependent variables of age and relative adiposity, the wide range of estradiol (E2) concentrations observed in study subjects, and a sex-specific effect of age on GH secretion (17) might have each played a role in the conclusions reached in the various studies.

In this study, we have defined gender-specific patterns of spontaneous and GHRH-stimulated GH secretion through intensive blood sampling and use of a super-sensitive GH assay in carefully selected groups of young men and women. In addition, we have investigated whether negative feedback by IGF-I on GH secretion is the same in men and women.

1. Abbreviations used in this paper: BPF, block-GH pulse frequency; BTI, block-total GH input; E2, estradiol; GH, growth hormone; GHRH, growth hormone–releasing hormone; IGF-I, insulin-like growth factor I; IGFBP-3, IGF binding protein 3; rhIGF-I, recombinant human IGF-I; P, progesterone; SRIH, somatostatin; TRH, thyrotropin-releasing hormone; TSH, thyroid-stimulating hormone; T3, triiodothyronine; T3RU, triiodothyronine resin uptake; T4, thyroxine.
Methods

Subjects. The research protocol was approved by the Institutional Review Board and the General Clinical Research Center Operating Subcommittee of the University of Michigan Hospitals. Written consent was obtained from each subject before their participation. Eight healthy men and eight healthy women were recruited. The two groups were of similar age (23.5 ± 2.9 vs. 25.2 ± 4.0 yr; mean ± SD; men vs. women) and body mass index (23.8 ± 0.9 vs. 22.9 ± 2.9 kg/m²). All had unremarkable medical histories and physical examinations. Measurements of renal, hepatic, and hematological function in all subjects were normal. None of the subjects were on any medications. Only women with regular spontaneous menstrual cycles were included. Women were studied during the early follicular phase of the cycle and menstrual status was further validated by measurement of serum estradiol (E2) and progesterone (P) concentrations.

Protocol. The subjects were admitted to the GCRC the evening before the actual study and antegrade intravenous cannulae were placed in forearm veins of each arm. The subjects were fed three meals each day served at 0700, 1200, and 1800 h, respectively. Caloric content was standardized as previously described (18). Lights were turned on at 0630 h and off at 2300 h. Daytime napping was not allowed.

Eight men were each studied twice, once while receiving normal saline (control) and once while receiving recombinant human IGF-I (rhIGF-I; Genentech, San Francisco, CA). Data on six of these men during the control saline infusion and on four men during the infusion of 10 μg/kg/h rhIGF-I have been previously reported (18). Two additional men were studied using saline and 10 μg/kg/h rhIGF-I. Both studies in the men were performed during the same admission as previously described (18). In brief, every 10 min, blood sampling was done for GH from 0800 h, day 1 until 1200 h, day 2. At 0800 h, day 2, an i.v. bolus of 50 μg thyrotropin-releasing hormone (TRH, Thyropinone; Abbot Laboratories, North Chicago, IL) was given. This was followed by administration of an i.v. bolus of 0.3 μg/kg/h GHRH-44 (Bachem California, Torrance, CA) at 1000 h. After completion of the GHRH test at 1200 h, an infusion of 10 μg/kg body weight/h rhIGF-I was begun and continued until 1200 h, day 4. Every 10 min, sampling for GH was again performed from 0800 h, day 3 until 1200 h, day 4. I.v. boluses of TRH and GHRH were given on day 4 in doses and times identical to that on day 2. For the women, a similar protocol was followed; however, the saline and IGF-I studies were performed in random order during the early follicular phase of different menstrual cycles. Care was exercised to match the timing of saline and rhIGF-I infusion to the same day of the menstrual cycle and women were admitted to the GCRC for study within 48 h of the onset of menses. For both men and women, plasma IGF-I was measured at 4-h intervals during the saline and IGF-I infusions. Plasma samples for total thyroxine (T4), T3 resin uptake (T3RU), and total triiodothyronine (T3) were obtained just before the TRH administration. Blood was sampled every 20 min for thyroid-stimulating hormone (TSH) from times 20 to +60 min with respect to the TRH bolus. Plasma E2, P, and testosterone (T) were measured in a sample made by pooling equal aliquots of plasma obtained every 6 h during the period of frequent blood sampling.

Assays. Plasma GH was measured in duplicate by a chemiluminescent assay (Nichols Institute Diagnostics, San Juan Capistrano, CA) that is specific for 22 KD GH as previously described (19). All samples from each particular subject were analyzed in the same assay and the detection limit of the assay was 0.01 μg/liter. Mean intraassay coefficient of variation (CV) was 9% between 0.01 and 0.1 μg/liter and 5% between 0.1 and 40 μg/liter. Interassay CV was 7% at 9 μg/liter. rhIGF-I at a concentration of 1,000 μg/liter did not interfere with the measurement of GH. Total IGF-I was measured after acid-ethanol extraction by an immunoradiometric assay (Diagnostics Systems Laboratories; Webster, TX) and IGF binding protein-3 (IGFBP-3) was measured using RIA kits (Diagnostics Systems Laboratories). All IGF-I measurements for a particular patient were performed in a single assay and the interassay and mean intraassay coefficients of variations were 6 and 5%, respectively. Plasma E2, P, and T were measured using commercial kits (Coat-A-Count, Diagnostics Products Corporation; Los Angeles, CA). All measurements of IGFBP-3, E2, P, and T were performed in single assays. Total T4, total T3, T3RU, and TSH assays were performed by the Ligand Laboratory of the University of Michigan Medical Center using standard methodologies.

Data analysis. Integrated GH concentration (in micrograms × minute per liter) was calculated as the area under the GH vs. time curve using the trapezoidal rule. This was performed for the time period 0800-0800 h during both saline and IGF-I infusions. The GH responses to GHRH and the TSH responses to TRH were defined as maximal increases in the hormone concentrations over the baseline.

Parameters of GH pulsatility were analyzed with regard to both concentration and secretion. Cluster analysis was used to define peaks in the plasma GH concentration profiles as previous described (18). An absolute nadir was defined as the minimum GH concentration over 24 h for individual subjects. Wave form–dependent deconvolution analysis (20) of the 24-h GH profiles was used to calculate GH half-life, GH pulse frequency, and GH secretion profile. The absolute GH nadir in each series was assumed to reflect basal GH secretion. Daily GH secretion (micrograms per 24 h) was calculated as the daily GH secretion (micrograms per liter) times the total volume of distribution (Lv), which was assumed to be 7.9% of body weight (21). GH concentrations and discrete parameters of pulsatile GH secretion were compared between men and women during saline and IGF-I treatments using two-tailed, two-sample t tests.

As a measure of uniformity of GH pulse amplitude, a SD for each subject’s GH pulses during the baseline study was calculated. GH pulse amplitudes were logarithmically transformed before analysis. A similar analysis was performed after logarithmic transformation of the 145 daily GH concentrations. To determine whether the daily GH secretory pattern differed between men and women, we divided the 24-h sampling period into eight 3-h blocks and calculated the block total GH input (BTI) and the block-GH pulse frequency (BPF) based on the estimates from deconvolution. These data were analyzed by repeated measures ANOVA. The percent of the day spent in actual active GH secretion was calculated from the deconvolution output.

<table>
<thead>
<tr>
<th>Parameter</th>
<th>Men (n = 8)</th>
<th>Women (n = 8)</th>
<th>P</th>
</tr>
</thead>
<tbody>
<tr>
<td>GH concentration analysis (cluster)</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>IGHC (μg × min/liter)</td>
<td>2884±661</td>
<td>3621±613</td>
<td>NS</td>
</tr>
<tr>
<td>Mean pulse amplitude (μg/liter)</td>
<td>5.2±0.8</td>
<td>3.8±0.7</td>
<td>NS</td>
</tr>
<tr>
<td>Median pulse amplitude (μg/liter)</td>
<td>1.3±0.3</td>
<td>1.5±0.2</td>
<td></td>
</tr>
<tr>
<td>Maximum pulse amplitude (μg/liter)</td>
<td>24.1±3.6</td>
<td>12.1±2.6</td>
<td>0.017</td>
</tr>
<tr>
<td>Mean valley (μg/liter)</td>
<td>0.13±0.02</td>
<td>0.28±0.03</td>
<td>0.001</td>
</tr>
<tr>
<td>Absolute nadir (μg/liter)</td>
<td>0.06±0.01</td>
<td>0.07±0.01</td>
<td>NS</td>
</tr>
<tr>
<td>Pulse frequency (per 24 h)</td>
<td>8.2±0.8</td>
<td>10.3±0.5</td>
<td>0.05</td>
</tr>
<tr>
<td>SD of pulse amplitudes (μg/liter)</td>
<td>7.98±1.10</td>
<td>5.31±1.04</td>
<td>0.001</td>
</tr>
<tr>
<td>GH secretory analysis (deconvolution)</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Daily GH secretion rate (μg/Lv per 24 h)</td>
<td>93.2±20.8</td>
<td>105.0±14.8</td>
<td>NS</td>
</tr>
<tr>
<td>Total GH secreted (μg/24 h)</td>
<td>561±130</td>
<td>514±88</td>
<td></td>
</tr>
<tr>
<td>Pulse frequency (per 24 h)</td>
<td>10.8±1.1</td>
<td>19.9±2.3</td>
<td>0.005</td>
</tr>
<tr>
<td>Total duration of pulsatile secretion (per 24 h)</td>
<td>24±2</td>
<td>42±4</td>
<td>0.003</td>
</tr>
<tr>
<td>GH half-life (min)</td>
<td>16.5±0.6</td>
<td>18.1±1.0</td>
<td>NS</td>
</tr>
</tbody>
</table>
Repeated measures ANOVA was performed to determine the effects of gender and of rhIGF-I on GH secretion and the potential interaction between these two factors. The effects of rhIGF-I on sex steroids, IGFBP-3, and thyroid hormones were analyzed by paired t tests. In all comparisons, data that were not normally distributed were logarithmically or square root transformed as appropriate before analyses. $P < 0.05$ was considered statistically significant. Data are presented as mean±SE.

**Results**

**Baseline GH secretion.** A comparison of 24-h GH concentration and secretion parameters between men and women during the control day is presented in Table I. Total daily GH secretion as measured by 24-h IGHC was similar between the two genders. Women had marginally more pulses ($P = 0.05$). There were no differences between men and women with regard to either mean or median pulse amplitude, but the maximum pulse amplitude was greater in men ($P = 0.017$). Although the nadirs for the 24-h baseline studies were similar in the two groups ($P = 0.5$), mean GH valley concentrations were twice as high in women ($P = 0.001$).

Similar to the results of the 24-h IGHC, the deconvolution analysis also demonstrated that total GH input was indistinguishable between the two groups ($P = 0.46$). Total daily GH production rate was also identical in the two groups ($P = 0.77$). Men had fewer GH secretory events per 24 hours than did women (10.8±1.1 vs. 19.9±2.3 secretory pulses/24 h; $P = 0.005$). Deconvolution-estimated GH half-life and mean secretory amplitude were similar in the two groups. The only significant association found by regression analysis of mean GH with plasma E2, P, T, and IGF-I was a positive correlation between GH and E2 in men ($r^2 = 0.63; P = 0.02$).

Although there were no gender-related effects on total GH secretion, there were clear differences in the pattern in which men and women secreted the hormone. In Fig. 1, representative GH concentration profiles of two men and two women are plotted on linear (top) and logarithmic (middle) scales. Decon-
volution-estimated GH secretory patterns are plotted in the bottom panel and the GH concentrations derived from these secretion estimates are overlaid on the measured concentration profiles. The mean (±SE) GH concentrations for the eight men and eight women are given in Fig. 2. The 24-h profiles in men were characterized by the presence of a dominant nocturnal pulse with much smaller pulses at other times of the day. In contrast, GH secretion in women was more continuous, with pulses of similar amplitude throughout the 24 h.

Several methods were used to quantify the differences in the pattern of GH secretion in men and women (Table I). As a measure of uniformity of pulse amplitude, a SD for each subject’s GH pulse amplitudes during the baseline study was calculated. GH pulse amplitudes were logarithmically transformed before analysis. This estimate of variation was greater in men than in women (7.98 vs. 5.31 μg/liter; P = 0.001). A similar analysis was performed after logarithmic transformation of the 145 daily GH concentrations. A higher SD for daily GH was found in men (5.22 vs. 4.03 μg/liter; P = 0.04). Fig. 3 shows a histogram of GH concentrations in these two groups. GH concentrations were either equal to or above the assay detection limit 98% of the time in men and 100% of the time in women. Although there was no difference between men and women in terms of absolute GH nadir, there was a difference in the frequency of biologically low GH concentrations. Reuten et al. (22) recently reported that subjects with severe organic GH deficiency had virtually all plasma GH concentrations, <0.5 μg/liter during 24-h sampling. Using this value as an estimate of the minimum GH concentration for bioactivity, only 35±4% of the spontaneous plasma GH measurements in men were above this value, whereas 53±6% of the concentrations in women exceeded this limit (P = 0.04).

Deconvolution was used to determine differences in GH secretion that would account for gender-specific GH concentration profiles. Fig. 4 (top) shows the BPF for men and women calculated as the deconvolution estimated pulse fre-
frequency in each of eight 3-h time blocks. There was both a gender effect \((P = 0.002)\) and a time effect \((P = 0.0001)\). Women had more pulses than men and the number of GH pulses was greatest for both genders between 2300 and 0200 h. There was also an interaction between the gender and time effects \((P = 0.003)\), indicating that the pattern of GH pulse frequency was different between the two groups, with pulse frequency more evenly distributed over the 24-h study period in women. The analysis of the deconvolution net input (BTI) within these same time blocks is given in the bottom panel. There was a gender effect \((P = 0.004)\) and a time effect \((P = 0.0001)\) as well as a gender–time interaction \((P = 0.0005)\) for BTI. The strong gender–time interaction was a result of a much greater time effect in men \((P = 0.0001)\) than in women \((P = 0.05)\). The GH input at each time block, except during the expected nocturnal augmentation of GH, was greater in women than in men, suggesting that women secrete GH more continuously throughout the day, whereas men have predominantly nocturnal GH secretion. This was further demonstrated by deconvolution analysis, which estimated that active secretion occurred 42±4% of the time in women but only 24±2% of the time in men \((P = 0.003)\).

**Effect of rhIGF-I on plasma IGF-I and IGFBP-3 concentrations and GH secretion.** Mean IGF-I concentrations for the six men and seven women who received 10 \(\mu g/\text{kg/h} \) rhIGF-I are shown in Fig. 5. At baseline, the IGF-I concentrations for the two groups were similar \((231±21 \text{ vs. } 262±25 \mu g/\text{liter}; P = 0.4)\). Approximately 8 h after the infusion was started, the concentration curves for men and women began to diverge and after 20 h a plateau in IGF-I concentration was reached for both sexes. Mean IGF-I concentrations during the 0800–0800-h blood sampling period were lower in men than in women \((1089±41 \text{ vs. } 1389±99 \mu g/\text{liter}; P = 0.02)\). Before the rhIGF-I infusion, plasma IGFBP-3 concentrations were identical in the two groups \((3.00±0.07 \text{ vs. } 3.02±0.24 \text{ mg/liter}; P = 0.9)\). Although GH secretion fell during rhIGF-I infusion, plasma IGFBP-3 actually increased in men \((P < 0.005)\) and there was a trend for higher levels in women.

The effects of rhIGF-I infusion on GH concentration in men (left) and women (right) is shown in Fig. 6 and discrete parameters of pulsatile GH secretion for the six men and seven women who received rhIGF-I are given in Table II. In both sexes, infusion of rhIGF-I suppressed pulsatile GH secretion and both total 24-h IGHC and pulse frequency were significantly lower during rhIGF-I treatment. The degree of suppression, however was gender specific and the same dose of rhIGF-I resulted in 84±3% suppression of the 24-h IGHC in men, but only 64±6% suppression in women \((P = 0.007)\). A comparison between the effect of rhIGF-I on the deconvolution output showed that suppression of GH secretion was similarly sexually dimorphic and that the degree of suppression on the calculated secretion was virtually identical to IGHC suppression. rhIGF-I had no effect on GH half-life as calculated by deconvolution.

**GH responses to GHRH (Fig. 7).** By ANOVA, there was no gender effect on the GH response to GHRH during the control study. There was, however an interaction between gender and treatment. The GH response to the GHRH bolus was significantly attenuated in men by the rhIGF-I infusion \((18.66±2.98 \text{ vs. } 3.88±0.74 \mu g/\text{liter}; P = 0.01)\), with a mean suppression of 75±9%. In contrast, rhIGF-I infusion had no effect on the women’s GH response to GHRH \((19.69±6.30 \text{ vs. } 24.22±8.14 \mu g/\text{liter}; P = 0.6)\). There was no correlation between the degree of suppression of spontaneous and GHRH stimulated GH secretion.

**Effects of rhIGF-I on thyroid hormones and sex steroids (Fig. 8, Table III).** By two-way ANOVA, there were no gender effects on T4, T3, T3RU, or TSH. Hence, in subsequent analyses, data from men and women were combined. rhIGF-I had no effect on T4 or T3RU, however it resulted in an increase in T3 \((101±7.8 \text{ vs. } 126.7±7.0 \text{ ng/dl}; P = 0.001)\), suppression of baseline TSH \((1.51±0.19 \text{ vs. } 0.64±0.09 \text{ mU/liter}; P < 0.0001)\) and suppression of the TSH increase after the TRH bolus \((6.83±0.55 \text{ vs. } 4.41±0.61 \text{ mU/liter}; P < 0.0001)\). The mean degree of suppression of the TSH response to TRH was 37±9% in men and 39±8% in women. rhIGF-I had no effect on the concentrations of sex steroids in men. In women, baseline E2 and P concentrations were appropriate for early follicular phase and, in comparison to the control study, plasma E2 and T concentrations were significantly higher during the

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rhIGF-I treatment. rhIGF-I infusion had no effect on P concentrations.

Discussion

This study demonstrates the existence of sexual dimorphism in the regulation of GH secretion. We show that gender-related differences are manifested by the patterns of GH secretion, rather than the total daily GH secretion. Moreover, we show that there are striking gender-related differences in the GH responses to negative feedback by IGF-I.

Several previous studies have suggested that GH secretion in humans is sexually dimorphic, as manifested by higher GH secretory rates in women (5, 6). In contrast to these earlier observations, we found that overall daily GH secretion rates were similar between men and women. A power analysis using integrated growth hormone concentration in men and women, where the effect size was 0.4 SD, suggests that 100 subjects per group would be needed to show a difference with 80% power. Thus, even if the difference between men and women is real, it is unlikely to be biologically meaningful. A potential explanation for the different conclusions between this and other studies was the age of subjects. In our study, all subjects were between 18 and 30 yr old. In contrast, in other studies (5, 6), the subjects were older and the age ranges were wider. Since the age-related fall in GH secretion is more pronounced in men than in women (17), inclusion of older men would decrease the estimate of mean daily GH secretion for the male group and might account for the previously reported gender difference.

Our conclusions are supported by a study from Ho et al. (4), which included men and women in an age group similar to that of our study. They reported that higher integrated GH concentrations in young women were determined by serum estradiol concentrations, not by gender. Since women in our
study were studied in the early follicular phase, a time when estradiol levels were comparable between the sexes, GH secretion was similar in both groups. Thus, when the study populations were tightly matched for age, relative adiposity, and E2 concentrations, the absolute amounts of GH secreted per day were indistinguishable between men and women.

The positive correlation between plasma E2 and GH secretion in men observed in this study was also consistent with data from Ho et al. (4). Plasma E2 concentrations were uniformly low in the early–follicular phase women and this homogeneity probably obscured any potential correlation between plasma GH and E2 concentrations in this group. In order to more conclusively derive a relationship between GH secretion and sex steroids, studies across the menstrual cycle might be required.

Although the total GH secretion rates were similar in both groups, there were gender-specific differences in the patterns of GH secretion. First, women had more GH pulses as detected both by cluster and by deconvolution analysis. Second, the interpulse GH levels in women were twice as high as in men, similar to the male vs. female difference found in rats (1). Third, the temporal organization of the GH pulsatility was different in men and women.

GH secretion in men was characterized by the predominance of large nocturnal pulses and relatively low GH output during the rest of the day. In contrast, women had much more uniform GH pulses throughout the day. This gender-based difference in GH secretion temporal architecture was demonstrated by frequency histograms. 35% of GH concentrations in men, but 54% of concentrations in women exceeded 0.5 μg/liter, an estimate for the lower limit of biologically meaningful GH. These differences in GH concentration were shown to originate from different GH secretory patterns. Deconvolution analysis demonstrated that women had more frequent GH secretory events, had more uniform GH secretion throughout the day and spent nearly twice as much time in active GH secretion than did men.

These differences in GH pulse pattern may subserve possible sex-specific effects of GH. In rodents, sexually dimorphic GH pulse patterns are important for the regulation of many of the metabolic effects of GH, including muscle and cartilage.

Table II. Comparison of Parameters of GH Secretion and IGFBP-3 in Men and Women during Saline and rhIGF-I Infusions

<table>
<thead>
<tr>
<th>Parameter</th>
<th>Men (n = 6)</th>
<th>IGF-I</th>
<th>Women (n = 7)</th>
<th>IGF-I</th>
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</thead>
<tbody>
<tr>
<td>Concentration analysis (cluster)</td>
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<tr>
<td>IGHC (μg × min/liter)</td>
<td>3296±824</td>
<td>678±240*</td>
<td>3770±688</td>
<td>1246±232*</td>
</tr>
<tr>
<td>Pulse frequency (per 24 h)</td>
<td>9.1±0.6</td>
<td>5.5±0.5*</td>
<td>10.4±0.6</td>
<td>7.4±0.9*</td>
</tr>
<tr>
<td>Secretion analysis (deconvolution)</td>
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<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Secretion (μg/liter 24 h)</td>
<td>103.4±26.9</td>
<td>19.6±7.5*</td>
<td>107.8±16.8</td>
<td>37.2±7.9*</td>
</tr>
<tr>
<td>Pulse frequency (per 24 h)</td>
<td>11.5±1.3</td>
<td>10.2±1.2</td>
<td>20.1±2.6</td>
<td>16.0±2.3</td>
</tr>
<tr>
<td>Half-life (min)</td>
<td>17.3±0.1</td>
<td>16.2±0.1</td>
<td>18.3±0.2</td>
<td>18.4±0.2</td>
</tr>
<tr>
<td>IGFBP-3 (mg/liter)</td>
<td>3.00±0.07</td>
<td>3.58±0.12*</td>
<td>3.02±0.24</td>
<td>3.53±0.27</td>
</tr>
</tbody>
</table>

* P < 0.005 saline vs. rhIGF-I. † P < 0.001 saline vs. rhIGF-I.
IGF-I mRNA (23), GH receptor mRNA (24), hepatic P450 isoenzymes (2), lipoproteins (25), and somatic growth (3, 26). In the case of GH regulation of hepatic CYP2C11 and CYP2C12, Waxman et al. (2) have shown that it is neither the total GH exposure nor the GH pulse amplitude but rather the interval between pulses that is important for determining which isoform is expressed. Limited data support a role for GH in the expression of hepatic CYP1A2 in humans (27). In addition, we have previously demonstrated that GH alters the expression of the N-demethylating P450 enzyme CYP3A4; these preliminary data suggested that GH pulse pattern, not simply total GH secretion was important for this effect (28).

Whether gender-specific GH pulse patterns subserve similar regulation to that found in rodents will require careful clinical investigations. However, the differences in GH secretion and concentration profiles described in our experiments mirror data from animal studies. This provides a physiologic mechanism to explain our previous observations on gender differences in human CYP3A4 expression (28).

We had previously demonstrated that continuous infusion of rhIGF-I to men inhibited pulsatile GH secretion and the GH response to a GHRH bolus. We have extended our initial observations by studying whether the same regulatory mechanism holds for women.

Table III. Plasma Concentrations of Sex Steroids and Thyroid Hormones in Men and Women during Saline and rhIGF-I Infusions

<table>
<thead>
<tr>
<th>Parameter</th>
<th>Men (n = 6)</th>
<th>Women (n = 7)</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>Saline</td>
<td>IGF-I</td>
</tr>
<tr>
<td>E2 (ng/liter)</td>
<td>45.3±10.5</td>
<td>49.6±11.8</td>
</tr>
<tr>
<td>Progesterone (µg/liter)</td>
<td>0.68±0.03</td>
<td>0.64±0.06</td>
</tr>
<tr>
<td>Testosterone (µg/liter)</td>
<td>5.89±0.62</td>
<td>5.85±0.46</td>
</tr>
<tr>
<td>T4 (µg/dl)</td>
<td>6.9±0.4</td>
<td>7.7±1.1</td>
</tr>
<tr>
<td>T3RU (%)</td>
<td>104±6</td>
<td>97±4</td>
</tr>
<tr>
<td>T3 (ng/dl)</td>
<td>115±6</td>
<td>151±7*</td>
</tr>
<tr>
<td>Baseline TSH (mU/liter)</td>
<td>1.43±0.33</td>
<td>0.56±0.09†</td>
</tr>
</tbody>
</table>

*P < 0.05 saline vs. rhIGF-I. †P < 0.01 saline vs. rhIGF-I. ‡P < 0.005 saline vs. rhIGF-I.

**Figure 6.** Comparison of plasma GH concentration profiles (mean±SE) in six men (left) and seven women (right) who received 10 µg/kg/h rhIGF-I i.v.
Sexual Dimorphism of Growth Hormone Regulation

Significantly less effective in women than in men in terms of inhibiting both spontaneous and GHRH-stimulated GH, despite higher plasma IGF-I concentrations during the rhIGF-I infusion. The higher IGF-I concentrations in women are consistent with less GH suppression in this group so that the total IGF-I during the infusion might have reflected the summation of the rhIGF-I and the residual endogenous IGF-I production. A gender-specific difference in free IGF-I concentrations is unlikely, because plasma concentrations of IGFBP-3, the major IGF-I binding protein, increased more in men than in women during rhIGF-I infusion, suggesting that free IGF-I levels might have been lower in men. It is conceivable that direct pituitary suppression by rhIGF-I (29) is less effective in women. Investigation into this possibility will require in vitro study of normal somatotrophs from men and women.

These data suggest that sexual dimorphism of GH secretion in humans relates to at least two distinct phenomena: different patterns of spontaneous GH secretion and different suppressibility of spontaneous and GHRH-stimulated GH secretion by rhIGF-I. The neuroendocrine mechanisms that are responsible for these differences have not been fully established. Direct hypophysial–portal sampling in several nonhuman species has suggested that GH pulses are the result of hypothalamic discharges of GHRH (30–32), whereas SRIH likely sets the pituitary responsiveness to GHRH (32). In rats, intermittent secretion of SRIH could explain the periodic responsiveness to GHRH and characteristic GH pulse pattern seen in males of this species (33). This hypothesis is supported by lower levels of hypothalamic SRIH mRNA in female rats (34). In addition, SRIH may be less inducible by negative feedback by GH in females (35). Measurement of GHRH or SRIH in hypothalamic–pituitary blood is not practical in humans and peripheral measurements of these peptides may not reflect hypothalamic secretion. Our previous studies in humans using a specific antagonist to the GHRH receptor (36, 37), as well as physiologic experiments demonstrating concordance between GHRH and GH secretion in sheep (31, 32) suggest that GHRH and SRIH in higher mammals likely play roles that are parallel to those found in rats.

If this is the case, the gender-related differences in the pat-

Figure 7. Plasma GH responses (mean±SE) to GHRH (0.33 μg/kg i.v.) in men (top) and women (bottom) during saline (○) and rhIGF-I (●) infusions.
tern of GH secretion can be mainly attributed to the lesser role of SRIH in women. This would explain more continuous GH secretion, higher interpulse GH concentrations and the occurrence of more uniform and more frequent GH pulses in women. Since part of the IGF-I feedback is likely mediated through an augmentation of SRIH secretion (18), the attenuated inhibition of GH secretion by rhIGF-I in women would also support this hypothesis.

rhIGF-I-mediated suppression of spontaneous, but not GHRH-stimulated, GH secretion sets women apart from men, in whom both parameters declined in parallel. The simplest explanation for this difference might be a more prominent suppressive effect of rhIGF-I on GHRH neurons in women. IGF-I negative feedback on GHRH has been described in vitro (38) and was recently suggested by in vivo data in rats (39). This feedback could be sexually dimorphic and our results are similar to studies in rats in which exogenous GH suppressed both spontaneous and GHRH-stimulated GH secretion in males but only GHRH-induced GH release in females (35, 40). More powerful effects of SRIH in men could have overshadowed rhIGF-I suppression of GHRH. Moreover, less SRIH effect but similarity in total spontaneous GH secretion in men and women would require a compensatory decrease in GHRH secretion in women. Indeed, lower levels of GHRH mRNA have been found in hypothalami of female rats (41); median eminence GHRH peptide content is lower in ewes than in rams (42). Taken together, these data suggest that systemic IGF-I inhibits and stimulates hypothalamic GHRH and SRIH, respectively, and that the relative importance of the feedback on GHRH or SRIH neurons is sexually divergent. These interpretations are predicated on a model in which GH secretion is governed solely by GHRH and SRIH. It is also possible that the endogenous ligand for the recently cloned GH secretagogue receptor (43) plays a role in the sexual differentiation of GH secretion. Our data may need to be reinterpreted once this ligand is identified and its role in GH physiology is defined.

Figure 8. Plasma TSH responses (mean±SE) to TRH (50 μg i.v.) in men (top) and women (bottom) during saline (○) and rhIGF-I (●) infusions.
Baseline plasma TSH concentrations and TRH-stimulated TSH concentrations were equally suppressed by rhIGF-I infusion in both sexes. We had previously attributed this to a rhIGF-I-induced increase in SRIH (18). Our current data indicate that this might also reflect an increase in T4-to-T3 conversion during the rhIGF-I infusion. In contrast to our results, Klinger et al. (44) found no increase in T3 during rhIGF-I treatment, despite TSH suppression similar to what we observed. Of interest, an increase in T3 has also been found during GH treatment (45). Given the concomitant fall in plasma GH in our study, the mechanism of T3 rise during GH treatment is probably an IGF-I-mediated 5'-deiodinase effect rather than a result of GH per se. Whether the TSH suppression is due to an increase in SRIH, T3, or both mechanisms is uncertain.

During the rhIGF-I infusion, plasma E2 and T concentrations were higher in women than concentrations obtained during the control infusion. Both studies were performed in the early follicular phase of the menstrual cycle. Increased intraovarian steroid production driven by the systemic rhIGF-I infusion is therefore possible. IGF-I is produced locally in the ovary and may play an important role in ovarian regulation (46). Ovarian IGF-I might be either independent of or in part under the control of systemic GH (46). Our data suggest the intriguing possibility that circulating IGF-I, in addition to locally produced IGF-I, might play a role in intraovarian physiology.

In summary, we have demonstrated the existence of sexual dimorphism of the regulation of GH secretion in humans. This was expressed at a central level and involved spontaneous GH pulsatility as well as the negative feedback by IGF-I. We attribute this difference primarily to the more prominent role of hypothalamic SRIH in men than in women. Additionally, our data suggest that inhibition of hypothalamic GHRH secretion by IGF-I might play an important role in the control of GH secretion in women.

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References


